

## INFORMATION DISCLOSURE CITATION

(Use several sheets if necessary)

ATTY. DOCKET NO.  
D0273 NP  
APPLICATION NO.  
10/648,593  
APPLICANT  
HUANG ET AL.  
FILING DATE  
AUGUST 26, 2003

Group

## U.S. PATENT DOCUMENTS

EXAMINER INITIALS		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE
85	AA	US5569588	10/29/96	Ashby, et al.			
	AB						
	AC						
	AD						
	AE						
	AF						
	AG						
	AH						
	AI						
	AJ						
	AK						
	AL						

## FOREIGN PATENT DOCUMENTS

		DOCUMENT NUMBER	DATE	OFFICE	CLASS	SUBCLASS	TRANSLATION	
							YES	NO
	AM						<input type="checkbox"/>	<input type="checkbox"/>
	AN						<input type="checkbox"/>	<input type="checkbox"/>
	AO						<input type="checkbox"/>	<input type="checkbox"/>
	AP						<input type="checkbox"/>	<input type="checkbox"/>
	AQ						<input type="checkbox"/>	<input type="checkbox"/>

## OTHER DOCUMENTS (Including Author, Title, Date, Pertinent pages, Etc.)

85	AR	Cockett, et al., "Applied genomics: integration of the technology within pharmaceutical research and development", Current Opinion in Biotech, Vol. 11, pp. 602-609 (2000)
	AS	Sonneveld, P., "Multidrug resistance in haematological malignancies", J. Internal Med., Vol. 247, pp. 521-534 (2000)
4	AT	Alizadeh, et al., "Distinct types of diffuse large B-cell lymphoma identified by gene expression profiling", Nature, Vol. 403, pp. 503-511 (2000)

EXAMINER

SHERIDAN SWOPE, PH.D.  
PRIMARY EXAMINER

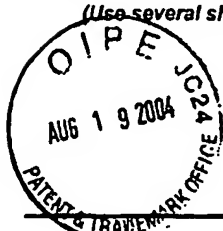
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2AA	Bittner, et al., "Molecular classification of cutaneous malignant melanoma by gene expression profiling", Nature, Vol. 406, pp. 536-540 (2000)
2AB	van't Veer, et al., "Gene expression profiling predicts clinical outcome of breast cancer", Nature, Vol. 15, pp. 530-536 (2002)
2AC	Khan, et al., "Classification and diagnostic prediction of cancers using gene expression profiling and artificial neural networks", Nature Medicine, Vol. 7, pp. 673-679 (2001)
2AD	Shipp, et al., "Diffuse large B-cell lymphoma outcome prediction by gene-expression profiling and supervised machine learning", Nature Medicine, Vol. 8, pp. 68-74 (2002)
2AE	Golub, et al., "Molecular Classification of Cancer: Class Discovery and Class Prediction by Gene Expression Monitoring", Science, Vol. 286, pp. 531-537 (1999)
2AF	Alon, et al., "Broad patterns of gene expression revealed by clustering analysis of tumor and normal colon tissues probed by oligonucleotide arrays", PNAS, Vol. 96, pp. 6745-6750 (1999)
2AG	West, et al., "Predicting the clinical status of human breast cancer by using gene expression profiles", PNAS, Vol. 98(20), pp. 11462-11467 (2001)
2AH	Sorlie, et al., "Gene expression patterns of breast carcinomas distinguish tumor subclasses with clinical implications", PNAS, Vol. 98(19), pp. 10869-10874 (2001)
2AI	Blanchard, et al., "Sequence to array: Probing the genome's secrets", Nature Biotechnology, Vol. 14, pp. 1649 (1996)
2AJ	Khan, et al., "Gene Expression Profiling of Alveolar Rhabdomyosarcoma with cDNA Microarrays", Cancer Res., Vol. 58, pp. 5009-5013 (1998)
2AK	Lockhart, et al., "Expression monitoring by hybridization to high-density oligonucleotide arrays", Nature Biotechnology, Vol. 14, pp. 1675-1680 (1996)
2AL	Freeman, et al., "Fundamentals of DNA Hybridization Arrays for Gene Expression Analysis", BioTechniques, Vol. 29, pp. 1042-1055 (2000)
2AM	Schena, et al., "Quantitative Monitoring of Gene Expression Patterns with a Complementary DNA Microarray", Science, Vol. 270, pp. 467-470 (1995)
2AN	

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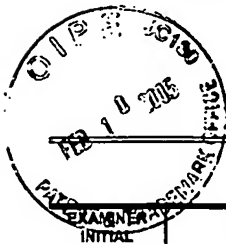
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	AR	Ellis, et al., "Down-regulation of Vascular Endothelial Growth Factor in a Human Colon Carcinoma Cell Line Transfected with an Antisense Expression Vector Specific for c-src", J. Biol. Chem., Vol. 273(2), pp. 1052-1057 (1998)
	AS	
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